

## SEQUENCE LISTING

<110> Nordlund, Henri Rainer et al.

<120> Avidin mutants

<130> BP110588

<160> 29

<170> PatentIn version 3.1

<210> 1

<211> 152

<212> PRT

<213> Gallus gallus

<400> 1

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Ala Leu Val Ala Pro Gly Leu Ser Ala Arg Lys Cys Ser Leu Thr Gly  
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Lys Trp Thr Asn Asp Leu Gly Ser Asn Met Thr Ile Gly Ala Val Asn  
35 40 45

Ser Arg Gly Glu Phe Thr Gly Thr Tyr Ile Thr Ala Val Thr Ala Thr  
50 55 60

Ser Asn Glu Ile Lys Glu Ser Pro Leu His Gly Thr Gln Asn Thr Ile  
65 70 75 80

Asn Lys Arg Thr Gln Pro Thr Phe Gly Phe Thr Val Asn Trp Lys Phe  
85 90 95

Ser Glu Ser Thr Thr Val Phe Thr Gly Gln Cys Phe Ile Asp Arg Asn  
100 105 110

Gly Lys Glu Val Leu Lys Thr Met Trp Leu Leu Arg Ser Ser Val Asn  
115 120 125

Asp Ile Gly Asp Asp Trp Lys Ala Thr Arg Val Gly Ile Asn Ile Phe  
130 135 140

Thr Arg Leu Arg Thr Gln Lys Glu  
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<211> 298

&lt;212&gt; PRT

&lt;213&gt; Gallus gallus

&lt;400&gt; 2

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1 5 10 15

Ala Leu Val Ala Pro Gly Leu Ser Ala Arg Lys Arg Thr Gln Pro Thr  
20 25 30

Phe Gly Phe Thr Val Asn Trp Lys Phe Ser Glu Ser Thr Thr Val Phe  
35 40 45

Thr Gly Gln Cys Phe Ile Asp Arg Asn Gly Lys Glu Val Leu Lys Thr  
50 55 60

Met Trp Leu Leu Arg Ser Ser Val Asn Asp Ile Gly Asp Asp Trp Lys  
65 70 75 80

Ala Thr Arg Val Gly Ile Asn Ile Phe Thr Arg Leu Arg Thr Gln Lys  
85 90 95

Glu Gly Gly Ser Gly Ser Ala Arg Lys Cys Ser Leu Thr Gly Lys  
100 105 110

Trp Thr Asn Asp Leu Gly Ser Asn Met Thr Ile Gly Ala Val Asn Ser  
115 120 125

Arg Gly Glu Phe Thr Gly Thr Tyr Ile Thr Ala Val Thr Ala Thr Ser  
130 135 140

Asn Glu Ile Lys Glu Ser Pro Leu His Gly Thr Gln Asn Thr Ile Asn  
145 150 155 160

Lys Ser Gly Gly Ser Thr Thr Val Phe Thr Gly Gln Cys Phe Ile Asp  
165 170 175

Arg Asn Gly Lys Glu Val Leu Lys Thr Met Trp Leu Leu Arg Ser Ser  
180 185 190

Val Asn Asp Ile Gly Asp Asp Trp Lys Ala Thr Arg Val Gly Ile Asn  
195 200 205

Ile Phe Thr Arg Leu Arg Thr Gln Lys Glu Gly Gly Ser Gly Gly Ser  
210 215 220

Ala Arg Lys Cys Ser Leu Thr Gly Lys Trp Thr Asn Asp Leu Gly Ser  
225 230 235 240

Asn Met Thr Ile Gly Ala Val Asn Ser Arg Gly Glu Phe Thr Gly Thr  
245 250 255

Tyr Ile Thr Ala Val Thr Ala Thr Ser Asn Glu Ile Lys Glu Ser Pro  
260 265 270

Leu His Gly Thr Gln Asn Thr Ile Asn Lys Arg Thr Gln Pro Thr Phe  
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Gly Phe Thr Val Asn Trp Lys Phe Ser Glu  
290 295

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cggctgtgat gtgggtgcct gtg 23

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<210> 24  
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<213> Gallus gallus

<400> 24

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20

25

30

Phe Gly Phe Thr Val Asn Trp Lys Phe Ser Glu Ser Thr Thr Val Phe  
35 40 45

Thr Gly Gln Cys Phe Ile Asp Arg Asn Gly Lys Glu Val Leu Lys Thr  
50 55 60

Met Trp Leu Leu Arg Ser Ser Val Asn Asp Ile Gly Asp Asp Trp Lys  
65 70 75 80

Ala Thr Arg Val Gly Ile Asn Ile Phe Thr Arg Leu Arg Thr Gln Lys  
85 90 95

Glu Gly Gly Ser Gly Gly Ser Ala Arg Lys Cys Ser Leu Thr Gly Lys  
100 105 110

Trp Thr Asn Asp Leu Gly Ser Asn Met Thr Ile Gly Ala Val Asn Ser  
115 120 125

Arg Gly Glu Phe Thr Gly Thr Tyr Ile Thr Ala Val Thr Ala Thr Ser  
130 135 140

Asn Glu Ile Lys Glu Ser Pro Leu His Gly Thr Gln Asn Thr Ile Asn  
145 150 155 160

Lys Ser Gly Gly Ser Thr Thr Val Phe Thr Gly Gln Cys Phe Ile Asp  
165 170 175

Arg Asn Gly Lys Glu Val Leu Lys Thr Met Trp Leu Leu Arg Ser Ser  
180 185 190

Val Asn Asp Ile Gly Asp Asp Trp Lys Ala Thr Arg Val Gly Ile Asn  
195 200 205

Ile Phe Thr Arg Leu Arg Thr Gln Lys Glu Gly Ser Gly Gly Ser  
210 215 220

Ala Arg Lys Cys Ser Leu Thr Gly Lys Trp Thr Asn Asp Leu Gly Ser  
225 230 235 240

Asn Met Thr Ile Gly Ala Val Asn Ser Arg Gly Glu Phe Thr Gly Thr  
245 250 255

Tyr Ile Thr Ala Val Thr Ala Thr Ser Asn Glu Ile Lys Glu Ser Pro  
260 265 270

Leu His Gly Thr Gln Asn Thr Ile Asn Lys Arg Thr Gln Pro Thr Phe  
275 280 285

Gly Phe Thr Val Asn Trp Lys Phe Ser Glu Gly Gly Ser Gly Ser Gly  
290 295 300

Ser Gly Ser Gly Ser Gly Arg Thr Gln Pro Thr Phe Gly Phe Thr Val  
305 310 315 320

Asn Trp Lys Phe Ser Glu Ser Thr Thr Val Phe Thr Gly Gln Cys Phe  
325 330 335

Ile Asp Arg Asn Gly Lys Glu Val Leu Lys Thr Met Trp Leu Leu Arg  
340 345 350

Ser Ser Val Asn Asp Ile Gly Asp Asp Trp Lys Ala Thr Arg Val Gly  
355 360 365

Ile Asn Ile Phe Thr Arg Leu Arg Thr Gln Lys Glu Gly Gly Ser Gly  
370 375 380

Gly Ser Ala Arg Lys Cys Ser Leu Thr Gly Lys Trp Thr Asn Asp Leu  
385 390 395 400

Gly Ser Asn Met Thr Ile Gly Ala Val Asn Ser Arg Gly Glu Phe Thr  
405 410 415

Gly Thr Tyr Ile Thr Ala Val Thr Ala Thr Ser Asn Glu Ile Lys Glu  
420 425 430

Ser Pro Leu His Gly Thr Gln Asn Thr Ile Asn Lys Ser Gly Gly Ser  
435 440 445

Thr Thr Val Phe Thr Gly Gln Cys Phe Ile Asp Arg Asn Gly Lys Glu  
450 455 460

Val Leu Lys Thr Met Trp Leu Leu Arg Ser Ser Val Asn Asp Ile Gly  
465 470 475 480

Asp Asp Trp Lys Ala Thr Arg Val Gly Ile Asn Ile Phe Thr Arg Leu  
485 490 495

Arg Thr Gln Lys Glu Gly Gly Ser Gly Ser Ala Arg Lys Cys Ser  
500 505 510

Leu Thr Gly Lys Trp Thr Asn Asp Leu Gly Ser Asn Met Thr Ile Gly  
 515 520 525

Ala Val Asn Ser Arg Gly Glu Phe Thr Gly Thr Tyr Ile Thr Ala Val  
 530 535 540

Thr Ala Thr Ser Asn Glu Ile Lys Glu Ser Pro Leu His Gly Thr Gln  
 545 550 555 560

Asn Thr Ile Asn Lys Arg Thr Gln Pro Thr Phe Gly Phe Thr Val Asn  
 565 570 575

Trp Lys Phe Ser Glu  
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<210> 25  
 <211> 1746  
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 <213> Gallus gallus  
 <221> DNA  
 <223> DNA sequence which codes for scAvd of SEQ ID NO 24

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 ttttcagagt ccaccactgt cttcacgggc cagtgcctca tagacaggaa tggaaaggag 180  
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 ggaggctccg ccagaaatgt ctcgctgact gggaaatgga ccaacgtatct gggctccaac 360  
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 aagtccggcg gatccaccac tgtcttcacg ggccagtgt tcatacgtacag gaatgggaag 540  
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 aacatgacca tcggggctgt gaacagcaga ggtgaattca caggcaccta catcacagcc 780  
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ggaggctccg gaggctccgc cagaaagtgc tcgctgactg ggaaatggac caacgatctg	1200
ggctccaaca tgaccatcgg ggctgtgaac agcagaggtg aattcacagg cacctacatc	1260
acagccgtaa cagccacatc aaatgagatc aaagagtcac cactgcatgg gacacaaaac	1320
accatcaaca agtccggcgg atccaccact gtcttacgg gccagtgctt catagacagg	1380
aatgggaagg aggtcctgaa gaccatgtgg ctgctgcgtt caagtgttaa tgacattggt	1440
gatgactgga aagctaccag ggtcggcatc aacatctca ctcgcctgcg cacacagaag	1500
gagggaggct ccggaggctc cgccagaaag tgctcgctga ctggaaatg gaccaacgat	1560
ctggctcca acatgaccat cgggctgtg aacagcagag gtgaattcac aggcacctac	1620
atcacagccg taacagccac atcaaatgag atcaaagagt caccactgca tggacaccaa	1680
aacaccatca acaagaggac ccagcccacc tttggctca ccgtcaatttgaagtttca	1740
gagtga	1746

<210> 26  
 <211> 897  
 <212> DNA  
 <213> Gallus gallus  
 <221> DNA  
 <223> DNA sequence which codes for dcAvd of SEQ ID 2

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ttttcagagt ccaccactgt ctacggc cagtgttca tagacaggaa tggaaaggag	180
gtcctgaaga ccatgtggct gtcgggtca agtgttaatg acattggta tgactggaaa	240
gctaccagg tcggcatcaa catttcaact cgcctgcgcac cacagaagga gggaggctcc	300
ggaggctccg ccagaaagtgc tgcgtgact gggaaatggac ccaacgatct gggctccaa	360
atgaccatcg gggctgtgaa cagcagaggtt gaattcacag gcacccatcat cacagccgt	420
acagccacat caaatgagat caaagagtca ccactgcgtg ggacacaaaaa caccatcaac	480
aagtccggcg gatccaccac tgtttcacg ggccagtgtt tcatacgacag gaatggaaag	540
gaggtcctga agaccatgtg gtcgtgcgg tcaagtgtta atgacattgg tgatgactgg	600
aaagctacca gggtcggcat caacatcttc actcgccctgc gcacacagaa ggagggaggc	660

tccggaggct ccgccagaaa gtgctcgctg actggaaat ggaccaacga tctggctcc	720
aacatgacca tcggggctgt gaacagcaga ggtgaattca caggcaccta catcacagcc	780
gtaacagcca catcaaatga gatcaaagag tcaccactgc atgggacaca aaacaccatc	840
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 <223> primer cp34\_C1

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<210> 28  
 <211> 290  
 <212> PRT  
 <213> Gallus gallus

<400> 28

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20	25
	30

Thr Val Asn Trp Lys Phe Ser Glu Ser Thr Thr Val Phe Thr Gly Gln	
35	40
	45

Cys Phe Ile Asp Arg Asn Gly Lys Glu Val Leu Lys Thr Met Trp Leu	
50	55
	60

Leu Arg Ser Ser Val Asn Asp Ile Gly Asp Asp Trp Lys Ala Thr Arg	
65	70
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	80

Val Gly Ile Asn Ile Phe Thr Arg Leu Arg Thr Gln Lys Glu Gly Gly	
85	90
	95

Ser Gly Gly Ser Ala Arg Lys Cys Ser Leu Thr Gly Lys Trp Thr Asn	
100	105
	110

Asp Leu Gly Ser Asn Met Thr Ile Gly Ala Val Asn Ser Arg Gly Glu	
115	120
	125

Phe Thr Gly Thr Tyr Ile Thr Ala Val Thr Ala Thr Ser Asn Glu Ile

130

135

140

Lys Glu Ser Pro Leu His Gly Thr Gln Asn Thr Ile Asn Lys Ser Gly  
 145 150 155 160

Gly Ser Lys Glu Ser Pro Leu His Gly Thr Gln Asn Thr Ile Asn Lys  
 165 170 175

Arg Thr Gln Pro Thr Phe Gly Phe Thr Val Asn Trp Lys Phe Ser Glu  
 180 185 190

Ser Thr Thr Val Phe Thr Gly Gln Cys Phe Ile Asp Arg Asn Gly Lys  
 195 200 205

Glu Val Leu Lys Thr Met Trp Leu Leu Arg Ser Ser Val Asn Asp Ile  
 210 215 220

Gly Asp Asp Trp Lys Ala Thr Arg Val Gly Ile Asn Ile Phe Thr Arg  
 225 230 235 240

Leu Arg Thr Gln Lys Glu Gly Ser Gly Gly Ser Ala Arg Lys Cys  
 245 250 255

Ser Leu Thr Gly Lys Trp Thr Asn Asp Leu Gly Ser Asn Met Thr Ile  
 260 265 270

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 275 280 285

Val Thr  
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<210> 29  
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 <212> DNA  
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